6114148

Syngp120mn

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 301 GTGCCCACCG ACCCCAACCC CCAGGAGGTG GAGCTCGTGA ACGTGACCGA
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 401 TCATCAGCCT GTGGGACCAG AGCCTGAAGC CCTGCGTGAA GCTGACCCCC
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Fig. 1A

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1501 GGCGGCGACA TGCGCGACAA CTGGAGATCT GAGCTGTACA AGTACAAGGT
1551 GGTGACGATC GAGCCCCTGG GCGTGGCCCC CACCAAGGCC AAGCGCCGCG
1601 TGGTGCAGCG CGAGAAGCGC TAAAGCGGCC GC
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(SEQ ID NO: 34)

Fig. 1B

Syngp160mn

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 101 AGGTGCACAA CGTGTGGGCC ACCCAGGCGT GCGTGCCCAC CGACCCCAAC
 151 CCCCAGGAGG TGGAGCTCGT GAACGTGACC GAGAACTTCA ACATGTGGAA
 201 GAACAACATG GTGGAGCAGA TGCATGAGGA CATCATCAGC CTGTGGGACC
 251 AGAGCCTGAA GCCCTGCGTG AAGCTGACCC CCCTGTGCGT GACCCTGAAC
 301 TGCACCGACC TGAGGAACAC CACCAACACC AACAACAGCA CCGCCAACAA
 351 CAACAGCAAC AGCGAGGGCA CCATCAAGGG CGGCGAGATG AAGAACTGCA
 401 GCTTCAACAT CACCACCAGC ATCCGCGACA AGATGCAGAA GGAGTACGCC
 451 CTGCTGTACA AGCTGGATAT CGTGAGCATC GACAACGACA GCACCAGCTA
 501 CCGCCTGATC TCCTGCAACA CCAGCGTGAT CACCCAGGCC TGCCCCAAGA
 551 TCAGCTTCGA GCCCATCCCC ATCCACTACT GCGCCCCCGC CGGCTTCGCC
 601 ATCCTGAAGT GCAACGACAA GAAGTTCAGC GGCAAGGGCA GCTGCAAGAA
 651 CGTGAGCACC GTGCAGTGCA CCCACGGCAT CCGGCCGGTG GTGAGCACCC
 701 AGCTCCTGCT GAACGGCAGC CTGGCCGAGG AGGAGGTGGT GATCCGCAGC
 751 GAGAACTTCA CCGACAACGC CAAGACCATC ATCGTGCACC TGAATGAGAG
 801 CGTGCAGATC AACTGCACGC GTCCCAACTA CAACAAGCGC AAGCGCATCC
 851 ACATCGGCCC CGGGCGCGCC TTCTACACCA CCAAGAACAT CATCGGCACC
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 951 GCGCCAGATC GTGAGCAAGC TGAAGGAGCA GTTCAAGAAC AAGACCATCG
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Fig. 1C

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1551 GAGCGGCATC GTGCAGCAGC AGAACAACCT CCTCCGCGCC ATCGAGGCCC
1601 AGCAGCATAT GCTCCAGCTC ACCGTGTGGG GCATCAAGCA GCTCCAGGCC
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1751 CCTCCTGGAG CAACAAGAGC CTGGACGACA TCTGGAACAA CATGACCTGG
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1951 CTGTGGTACA TCAAAATCTT CATCATGATT GTGGGCGGCC TGGTGGGCCT
2001 CCGCATCGTG TTCGCCGTGC TGAGCATCGT GAACCGCGTG CGCCAGGGCT
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2151 CGGCAGGCTC GTGCACGGCT TCCTGGCGAT CATCTGGGTC GACCTCCGCA
2201 GCCTGTTCCT GTTCAGCTAC CACCACCGCG ACCTGCTGCT GATCGCCGCC
2251 CGCATCGTGG AACTCCTAGG CCGCCGCGGC TGGGAGGTGC TGAAGTACTG
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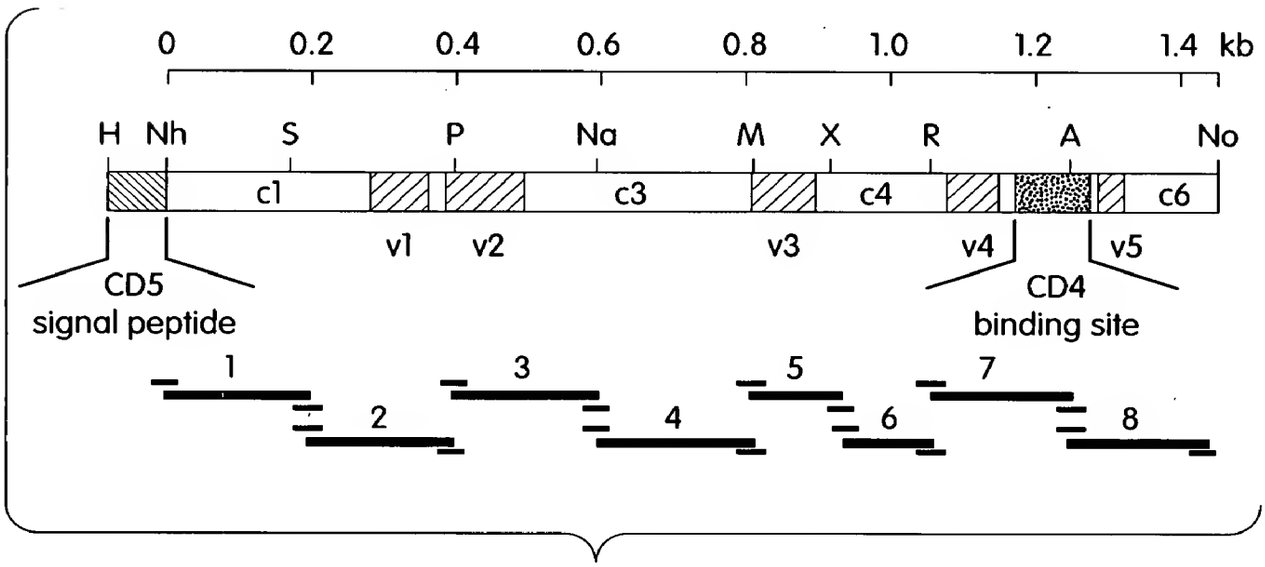
(SEQ ID NO: 35)

Fig. 1D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c6
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Fig. 2

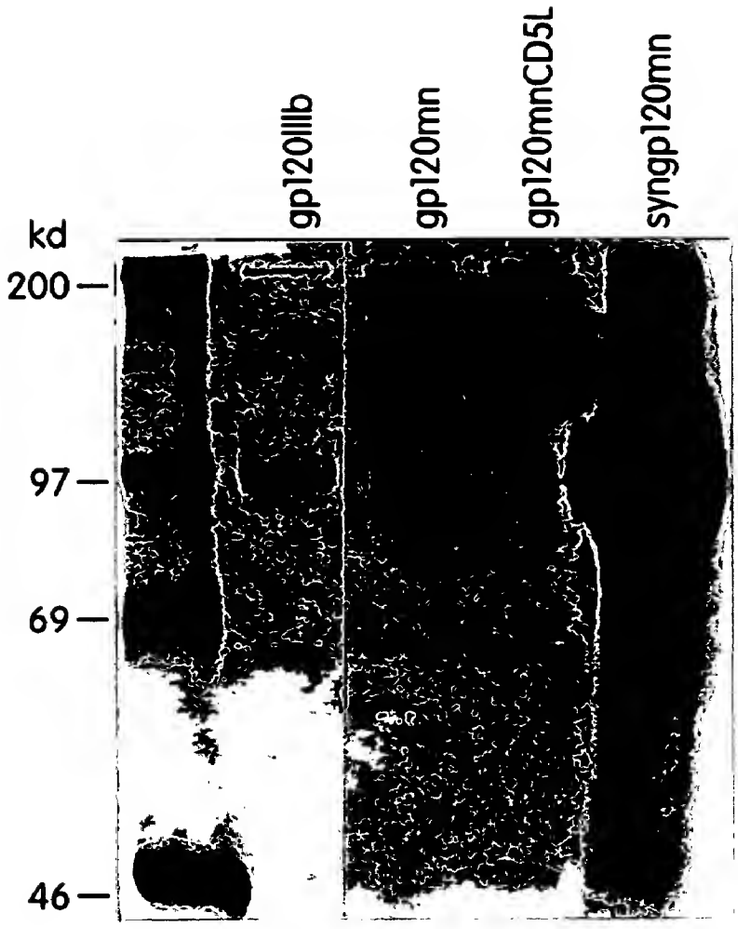

Fig. 3

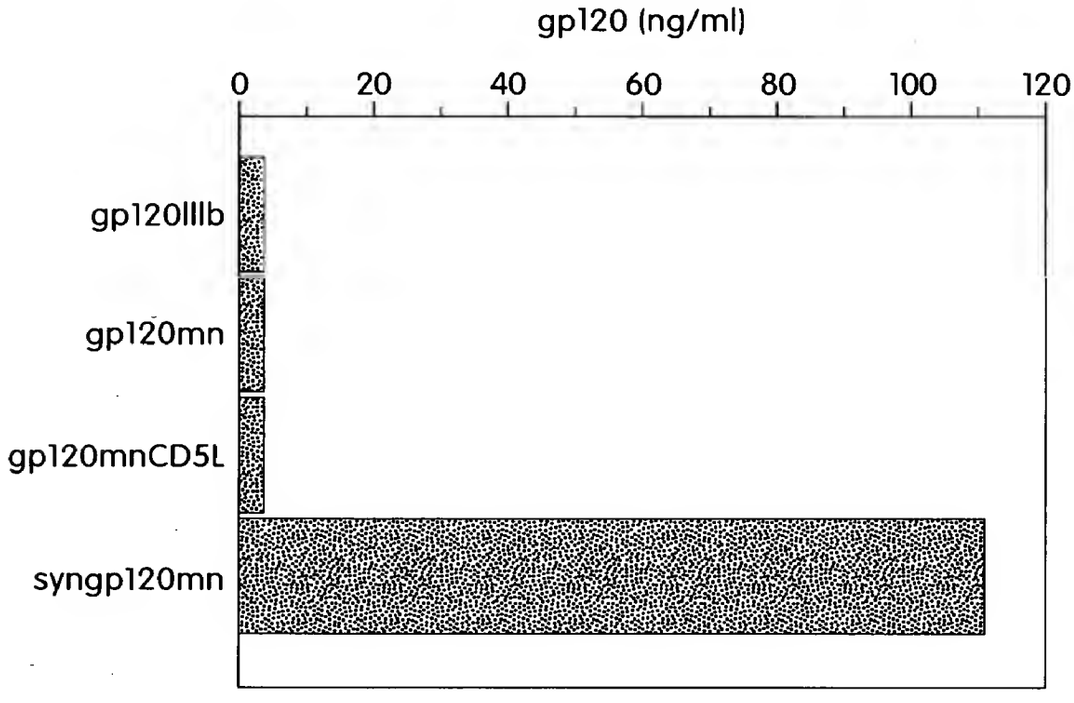
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Fig. 4

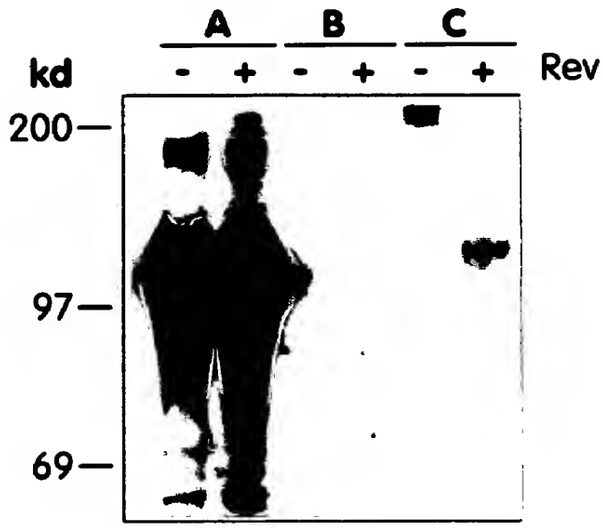

Fig. 5A

Fig. 5B

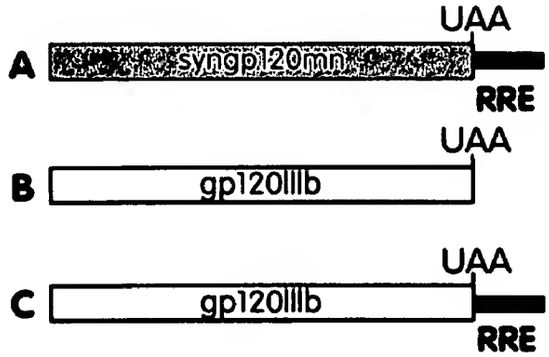

Fig. 5C

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      M   N   P   V   I   S   I   T   L   L   L   S   V   L   Q   M   S   R   G   Q

env  aga gta ata agt tta aca gca tgt tta gta aat caa aat ttg aga tta gat tgt aga cat 120
wt   agg gtg atc agc ctg aca gcc tgc ctg gtg aac cag aac ctt cga ctg gac tgc cgt cat
      R   V   I   S   L   T   A   C   L   V   N   Q   N   L   R   L   D   C   R   H

env  gaa aat aat aca aat ttg cca ata caa cat gaa ttt tca tta acg cgt gaa aaa aaa aaa 180
wt   gag aat aac acc aac ttg ccc atc cag cat gag ttc agc ctg acc cga gag aag aag aag
      E   N   N   T   N   L   P   I   Q   H   E   F   S   L   T   R   E   K   K   K

env  cat gta tta agt gga aca tta gga gta cca gaa cat aca tat aga agt aga gta aat ttg 240
wt   cac gtg ctg tca ggc acc ctg ggg gtt ccc gag cac act tac cgc tcc cgc gtc aac ctt
      H   V   L   S   G   T   L   G   V   P   E   H   T   Y   R   S   R   V   N   L

env  ttt agt gat aga ttc ata aaa gta tta aca tta gca aat ttt aca aca aaa gat gaa gga 300
wt   ttc agt gac cgc ttt atc aag gtc ctt act cta gcc aac ttc acc acc aag gat gag ggc
      F   S   D   R   F   I   K   V   L   T   L   A   N   F   T   T   K   D   E   G

env  gat tat atg tgt gag ctc aga gta agt gga caa aat cca aca agt agt aat aaa aca ata 360
wt   gac tac atg tgt gaa ctt cga gtc tcg ggc cag aat ccc aca agc tcc aat aaa act atc
      D   Y   M   C   E   L   R   V   S   G   Q   N   P   T   S   S   N   K   T   I

env  aat gta ata aga gat aaa tta gta aaa tgt gga gga ata agt tta tta gta caa aat aca 420
wt   aat gtg atc aga gac aag ctg gtc aag tgt ggt ggc ata agc ctg ctg gtt caa aac act
      N   V   I   R   D   K   L   V   K   C   G   G   I   S   L   L   V   Q   N   T

env  agt tgg tta tta tta tta tta tta agt tta agt ttt tta caa gca aca gat ttt ata agt 480
wt   tcc tgg ctg ctg ctg ctc ctg ctt tcc ctc tcc ttc ctc caa gcc acg gac ttc att tct
      S   W   L   L   L   L   L   L   S   L   S   F   L   Q   A   T   D   F   I   S

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wt   ctg tga            (SEQ ID NO: 37)
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Fig. 6

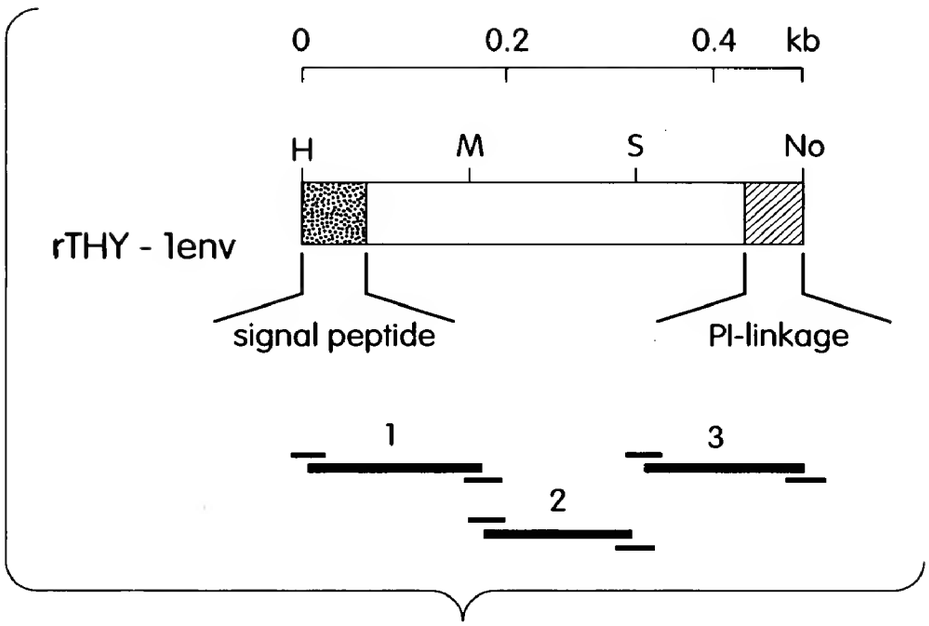

Fig. 7

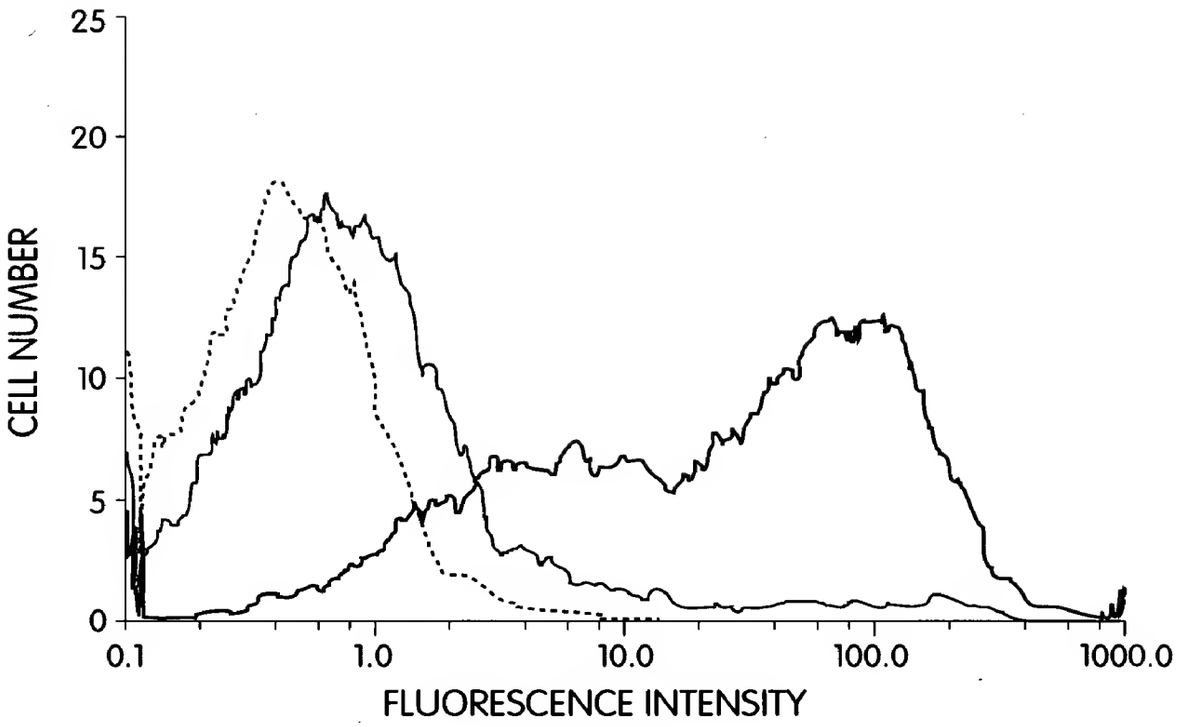
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Fig. 8

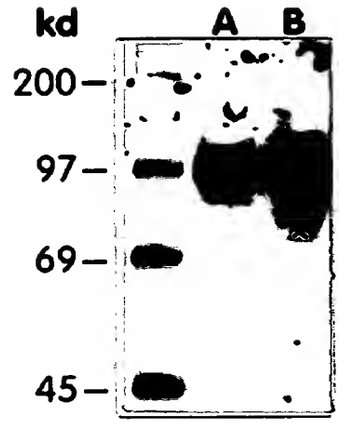

Fig. 9A

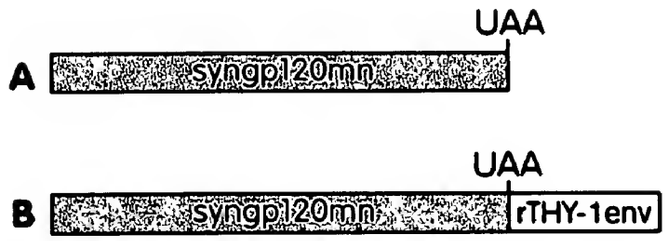

Fig. 9B

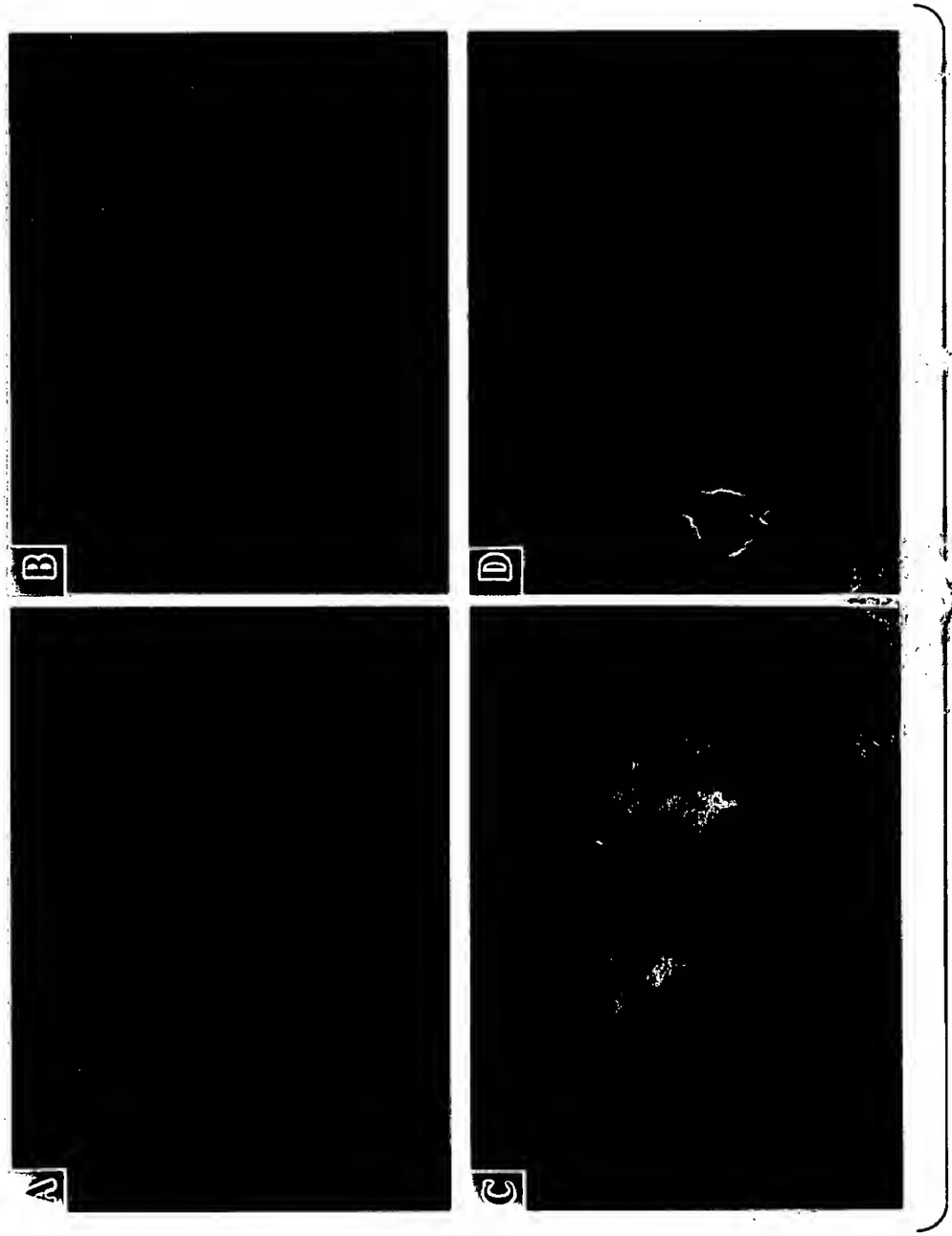

Fig. 10

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151 AAGCTGACCC TGAAGTTCAT CTGCACCACC GGCAAGCTGC CCGTGCCCTG
201 GCCCACCCTC GTGACCACCT TCAGCTACGG CGTGCAGTGC TTCAGCCGCT
251 ACCCCGACCA CATGAAGCAG CACGACTTCT TCAAGTCCGC CATGCCCGAA
301 GGCTACGTCC AGGAGCGCAC CATCTTCTTC AAGGACGACG GCAACTACAA
351 GACCCGCGCC GAGGTGAAGT TCGAGGGCGA CACCCTGGTG AACCGCATCG
401 AGCTGAAGGG CATCGACTTC AAGGAGGACG GCAACATCCT GGGGCACAAG
451 CTGGAGTACA ACTACAACAG CCACAACGTC TATATCATGG CCGACAAGCA
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551 GCAGCGTGCA GCTCGCCGAC CACTACCAGC AGAACACCCC CATCGGCGAC
601 GGCCCCGTGC TGCTGCCCGA CAACCACTAC CTGAGCACCC AGTCCGCCCT
651 GAGCAAAGAC CCCAACGAGA AGCGCGATCA CATGGTCCTG CTGGAGTTCG
701 TGACCGCCGC CGGGATCACT CACGGCATGG ACGAGCTGTA CAAGTAAAGC
751 GGCCGCGGAT CC
```

Fig. 11

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  51  CTGCTGGGGA TGCTGGTCGC TTCCGTGCTA GCCGCCACCA GAAGATACTA
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 251  TCACCTTTTC AACATCGCTA AGCCAAGGCC ACCCTGGATG GGTCTGCTAG
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 351  AACATGGCTT CCCATCCTGT CAGTCTTCAT GCTGTTGGTG TATCCTACTG
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Fig. 12A

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4201  CCAGCATGTA TGTGAAGGAG TTCCTCATCT CCAGCAGTCA AGATGGCCAT
4251  CAGTGGACTC TCTTTTTTCA GAATGGCAAA GTAAAGGTTT TTCAGGGAAA
4301  TCAAGACTCC TTCACACCTG TGGTGAACTC TCTAGACCCA CCGTTACTGA
4351  CTCGCTACCT TCGAATTCAC CCCCAGAGTT GGGTGCACCA GATTGCCCTG
4401  AGGATGGAGG TTCTGGGCTG CGAGGCACAG GACCTCTACT GAGGGTGGCC
4451  ACTGCAGCAC CTGCCACTGC CGTCACCTCT CCCTCCTCAG CTCCAGGGCA
4501  GTGTCCCTCC CTGGCTTGCC TTCTACCTTT GTGCTAAATC CTAGCAGACA
4551  CTGCCTTGAA GCCTCCTGAA TTAACTATCA TCAGTCCTGC ATTTCTTTGG
4601  TGGGGGGCCA GGAGGGTGCA TCCAATTTAA CTTAACTCTT ACCGTCGACC
4651  TGCAGGCCCA ACGCGGCCGC
```

Fig. 12B

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   1  AAGCTTAAAC CATGCCCATG GGGTCTCTGC AACCGCTGGC CACCTTGTAC
  51  CTGCTGGGGA TGCTGGTCGC TTCCGTGCTA GCCGCCACCC GCCGCTACTA
 101  CCTGGGCGCC GTGGAGCTGT CCTGGGACTA CATGCAGAGC GACCTGGGCG
 151  AGCTCCCCGT GGACGCCCGC TTCCCCCCCC GCGTGCCCAA GAGCTTCCCC
 201  TTCAACACCA GCGTGGTGTA CAAGAAAACC CTGTTCGTGG AGTTCACCGA
 251  CCACCTGTTC AACATTGCCA AGCCGCGCCC CCCCTGGATG GGCCTGCTGG
 301  GCCCCACCAT CCAGGCCGAG GTGTACGACA CCGTGGTGAT CACCCTGAAG
 351  AACATGGCCA GCCACCCCGT CAGCCTGCAC GCCGTGGGCG TGAGCTACTG
 401  GAAGGCCAGC GAGGGCGCCG AGTACGACGA CCAGACGTCC CAGCGCGAGA
 451  AGGAGGACGA CAAGGTGTTC CCGGGGGGGA GCCACACCTA CGTGTGGCAG
 501  GTGCTTAAGG AGAACGGCCC TATGGCCAGC GACCCCCTGT GCCTGACCTA
 551  CAGCTACCTG AGCCACGTGG ACCTGGTGAA GGATCTGAAC AGCGGGCTGA
 601  TCGGCGCCCT GCTGGTGTGT CGCGAGGGCA GCCTGGCCAA GGAGAAAACC
 651  CAGACCCTGC ACAAGTTCAT CCTGCTGTTC GCCGTGTTCG ACGAGGGGAA
 701  GAGCTGGCAC AGCGAGACTA AGAACAGCCT GATGCAGGAC CGCGACGCCG
 751  CCAGCGCCCG CGCCTGGCCC AAGATGCACA CCGTTAACGG CTACGTGAAC
 801  CGCAGCCTGC CCGGCCTGAT CGGCTGCCAC CGCAAGAGCG TGTACTGGCA
 851  CGTCATCGGC ATGGGCACCA CCCCTGAGGT GCACAGCATC TTCCTGGAGG
 901  GCCACACCTT CCTGGTGCGC AACCACCGCC AGGCCAGCCT GGAGATCAGC
 951  CCCATCACCT TCCTGACTGC CCAGACCCTG CTGATGGACC TAGGCCAGTT
1001  CCTGCTGTTC TGCCACATCA GCAGCCACCA GCACGACGGC ATGGAGGCTT
1051  ACGTGAAGGT GGACAGCTGC CCCGAGGAGC CCCAGCTGCG CATGAAGAAC
1101  AACGAGGAGG CCGAGGACTA CGACGACGAC CTGACCGACA GCGAGATGGA
1151  TGTCGTACGC TTCGACGACG ACAACAGCCC CAGCTTCATC CAGATCCGCA
1201  GCGTGGCCAA GAAGCACCCT AAGACCTGGG TGCACTACAT CGCCGCCGAG
1251  GAGGAGGACT GGGACTACGC CCCGCTAGTA CTGGCCCCCG ACGACCGCAG
1301  CTACAAGAGC CAGTACCTGA ACAACGGCCC CCAGCGCATC GGCCGCAAGT
1351  ACAAGAAGGT GCGCTTCATG GCCTACACCG ACGAGACTTT CAAGACCCGC
1401  GAGGCCATCC AGCACGAGTC CGGCATCCTC GGCCCCCTGC TGTACGGCGA
1451  GGTGGGCGAC ACCCTGCTGA TCATCTTCAA GAACCAGGCC AGCAGGCCCT
1501  ACAACATCTA CCCCCACGGC ATCACCGACG TGCGCCCCCT GTACAGCCGC
1551  CGCCTGCCCA AGGGCGTGAA GCACCTGAAG GACTTCCCCA TCCTGCCCGG
1601  CGAGATCTTC AAGTACAAGT GGACCGTGAC CGTGGAGGAC GGCCCCACCA
1651  AGAGCGACCC CCGCTGCCTG ACCCGCTACT ACAGCAGCTT CGTGAACATG
1701  GAGCGCGACC TGGCCTCCGG ACTGATCGGC CCCCTGCTGA TCTGCTACAA
1751  GGAGAGCGTG GACCAGCGCG GCAACCAGAT CATGAGCGAC AAGCGCAACG
1801  TGATCCTGTT CAGCGTGTTC GACGAGAACC GCAGCTGGTA TCTGACCGAG
1851  AACATCCAGC GCTTCCTGCC CAACCCCGCT GGCGTGCAGC TGGAAGATCC
1901  CGAGTTCCAG GCCAGCAACA TCATGCACAG CATCAACGGC TACGTGTTCG
1951  ACAGCCTGCA GCTGAGCGTG TGCCTGCATG AGGTGGCCTA CTGGTACATC
2001  CTGAGCATCG GCGCCCAGAC CGACTTCCTG AGCGTGTTCT TCTCCGGGTA
2051  TACCTTCAAG CACAAGATGG TGTACGAGGA CACCCTGACC CTGTTCCCCT
2101  TCTCCGGCGA GACTGTGTTC ATGTCTATGG AGAACCCCGG CCTGTGGATT
2151  CTGGGCTGCC ACAACAGCGA CTTCCGCAAC CGCGGCATGA CTGCCCTGCT
2201  GAAAGTCTCC AGCTGCGACA AGAACACCGG CGACTACTAC GAGGACAGCT
2251  ACGAGGACAT CTCCGCCTAC CTGCTGTCCA AGAACAACGC CATCGAGCCC
2301  CGCTCCTTCT CCCAAAACTC CCGCCACCCC AGCACGCGTC AGAAGCAGTT
2351  CAACGCCACC CCCCCCGTGC TGAAGCGCCA CCAGCGCGAG ATCACCCGCA
2401  CCACCCTGCA AAGCGACCAG GAGGAGATCG ACTACGACGA CACCATCAGC
2451  GTGGAGATGA AGAAGGAGGA CTTCGACATC TACGACGAGG ACGAGAACCA
2501  GAGCCCCCGC TCCTTCCAAA AGAAAACCCG CCACTACTTC ATCGCCGCCG
2551  TGGAGCGCCT GTGGGACTAC GGCATGAGCA GCAGCCCCCA CGTCCTGCGC
2601  AACCGCGCCC AGAGCGGCAG CGTGCCCCAG TTCAAGAAGG TGGTGTTCCA
2651  GGAGTTCACC GACGGCAGCT TCACCCAGCC CCTGTACCGC GGCGAGCTGA
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Fig. 13A

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2701  ACGAGCACCT GGGCCTGCTC GGCCCCTACA TCCGCGCCGA GGTGGAGGAC
2751  AACATCATGG TGACCTTCCG CAACCAAGCC TCCCGGCCCT ACTCCTTCTA
2801  CTCCTCCCTG ATCAGCTACG AGGAGGACCA GCGCCAGGGC GCCGAGCCCC
2851  GCAAGAACTT CGTGAAGCCC AACGAGACTA AGACCTACTT CTGGAAGGTG
2901  CAGCACCACA TGGCCCCCAC CAAGGACGAG TTCGACTGCA AGGCCTGGGC
2951  CTACTTCAGC GACGTGGACC TGGAGAAGGA CGTGCACAGC GGCCTGATCG
3001  GCCCCCTGCT GGTGTGCCAC ACCAACACCC TGAACCCCCC CCACGGGAGG
3051  CAGGTGACTG TGCAGGAATT TGCCCTGTTC TTCACCATCT TCGACGAGAC
3101  TAAGAGCTGG TACTTCACCG AGAACATGGA GCGCAACTGC CGCGCCCCCT
3151  GCAACATCCA GATGGAAGAT CCCACCTTCA AGGAGAACTA CCGCTTCCAC
3201  GCCATCAACG GCTACATCAT GGACACCCTG CCCGGCCTGG TGATGGCCCA
3251  GGACCAGCGC ATCCGCTGGT ACCTGCTGTC TATGGGCAGC AACGAGAACA
3301  TCCACAGCAT CCACTTCAGC GGCCACGTTT TCACCGTGCG CAAGAAGGAG
3351  GAGTACAAGA TGGCCCTGTA CAACCTGTAC CCCGGCGTGT TCGAGACTGT
3401  GGAGATGCTG CCCAGCAAGG CCGGGATCTG GCGCGTGGAG TGCCTGATCG
3451  GCGAGCACCT GCACGCCGGC ATGAGCACCC TGTTCCTGGT GTACAGCAAC
3501  AAGTGCCAGA CCCCCCTGGG CATGGCCAGC GGCCACATCC GCGACTTCCA
3551  GATCACCGCC AGCGGCCAGT ACGGCCAGTG GGCTCCCAAG CTGGCCCGCC
3601  TGCACTACAG CGGCAGCATC AACGCCTGGT CGACCAAGGA GCCCTTCTCC
3651  TGGATCAAGG TGGACCTGCT GGCCCCCATG ATCATCCACG GCATCAAGAC
3701  CCAGGGCGCC CGCCAGAAGT TCAGCAGCCT GTACATCAGC CAGTTCATCA
3751  TCATGTACTC TCTAGACGGC AAGAAGTGGC AGACCTACCG CGGCAACAGC
3801  ACCGGCACCC TGATGGTGTT CTTCGGCAAC GTGGACAGCA GCGGCATCAA
3851  GCACAACATC TTCAACCCCC CCATCATCGC CCGCTACATC CGCCTGCACC
3901  CCACCCACTA CAGCATCCGC AGCACCCTGC GCATGGAGCT GATGGGCTGC
3951  GACCTGAACA GCTGCAGCAT GCCCCTGGGC ATGGAGAGCA AGGCCATCAG
4001  CGACGCCCAG ATCACCGCCT CCAGCTACTT CACCAACATG TTCGCCACCT
4051  GGAGCCCCAG CAAGGCCCGC CTGCACCTGC AGGGCCGCAG CAACGCCTGG
4101  CGCCCCCAGG TGAACAACCC CAAGGAGTGG CTGCAGGTGG ACTTCCAGAA
4151  AACCATGAAG GTGACTGGCG TGACCACCCA GGGCGTCAAG AGCCTGCTGA
4201  CCAGCATGTA CGTGAAGGAG TTCCTGATCA GCAGCAGCCA GGACGGCCAC
4251  CAGTGGACCC TGTTCTTCCA AAACGGCAAG GTGAAGGTGT TCCAGGGCAA
4301  CCAGGACAGC TTCACACCGG TCGTGAACAG CCTGGACCCC CCCCTGCTGA
4351  CCCGCTACCT GCGCATCCAC CCCCAGAGCT GGGTGCACCA GATCGCCCTG
4401  CGCATGGAGG TGCTGGGCTG CGAGGCCCAG GACCTGTACT GAAGCGGCCG
4451  C
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Fig. 13B